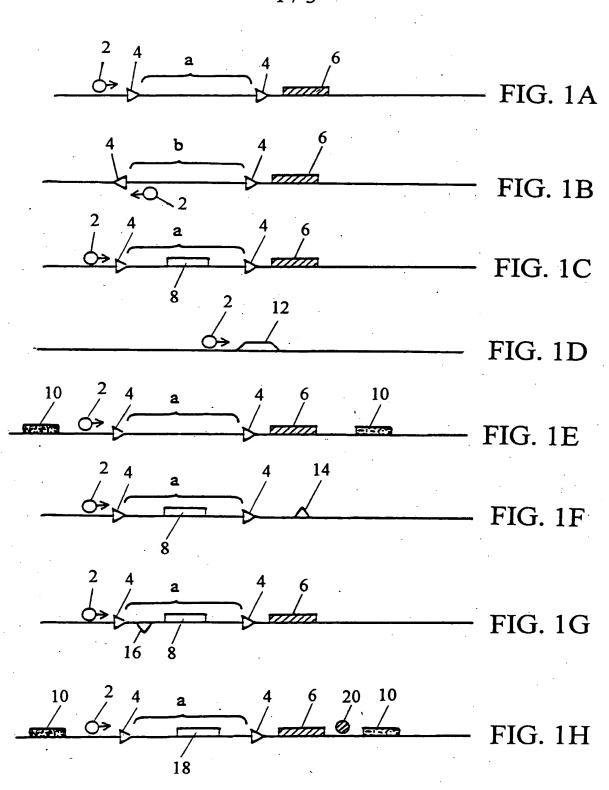
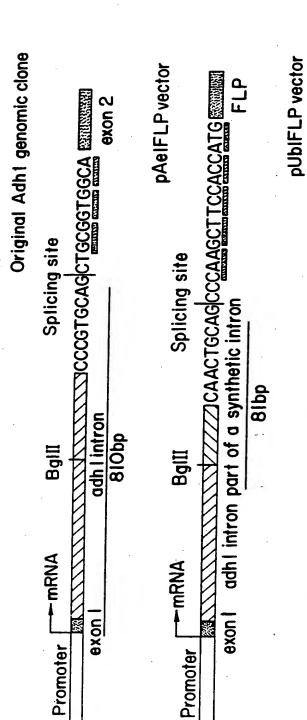
1/5



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FIG. 3



CGAAATCGATA AGCTTCCACCATG EXTREMENT □ CTTCTGCAG GTCGACTCTAGGATCCCCGGGTT Splicing site Ubi intron lolobp -mRNA BgIII exon | Promoter

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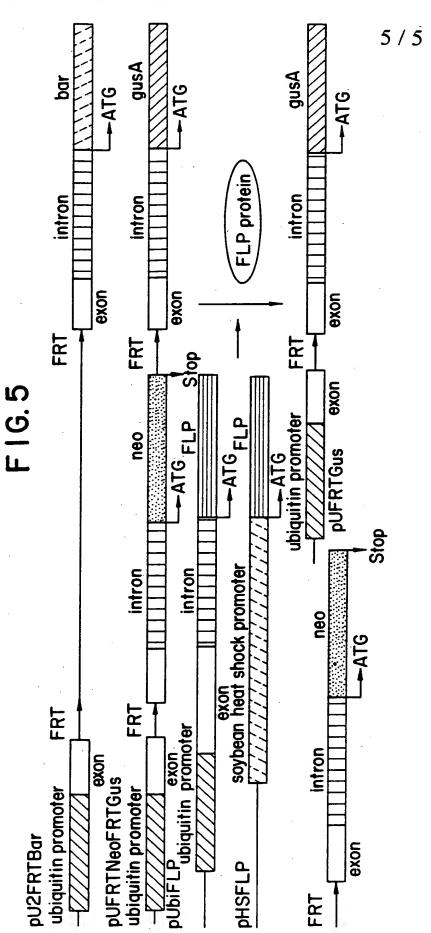
4/5 pU2FRTGrev. **pUFRTmG** pUbiGUS **pU2FRTG PUFRTG pUFRTG** GUS coding sequence poly A site poly A site poly A site FIG.4A | GATCAGAAGTTCCTATACTTTCTAGAGAATAGGAACTTCGGAATAGGAACTTCT GUS **/2 FRT** intron GUS coding sequence GUS coding sequence GATCAGTTCCTATACTTTCTAGAGAATAGGAACTTCGGAAT I.3kb pNEOBGAI ranslation start translation start fragment intron **1/2 FRT** exon exon L L FRT FRT Xbal XbaI defective FRTs 1.3kb pNEOBGAL exon Ubiquitin FRT promoter fragment fragment FRT Xbai IAGATCT promoter training training | mRNA exon Bgill exon exon Ubiquitin promoter F16.4B

<u>nokasack azasa</u>

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